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Last updated by author(s):	Dec 10, 2019

Reporting Summary

Nature Research wishes to improve the reproducibility of the work that we publish. This form provides structure for consistency and transparency in reporting. For further information on Nature Research policies, seeAuthors & Referees and theEditorial Policy Checklist.

Statistics	
For all statistical analys	es, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.
n/a Confirmed	
The exact sam	nple size (n) for each experimental group/condition, given as a discrete number and unit of measurement
A statement of	on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly
	test(s) used AND whether they are one- or two-sided ests should be described solely by name; describe more complex techniques in the Methods section.
A description	of all covariates tested
A description	of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons
	ion of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)
	thesis testing, the test statistic (e.g. F , t , r) with confidence intervals, effect sizes, degrees of freedom and P value noted is exact values whenever suitable.
For Bayesian a	analysis, information on the choice of priors and Markov chain Monte Carlo settings
For hierarchic	al and complex designs, identification of the appropriate level for tests and full reporting of outcomes
Estimates of e	effect sizes (e.g. Cohen's <i>d</i> , Pearson's <i>r</i>), indicating how they were calculated
1	Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.
Software and o	ode:
Policy information abo	ut <u>availability of computer code</u>
Data collection	PatchMaster (HEKA)
Data analysis	pClamp 10 (Molecular Devices); Sigmaplot (SPSS); MATLAB (Mathworks); Pymol (Schrodinger);
	om algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors/reviewers. deposition in a community repository (e.g. GitHub). See the Nature Research guidelines for submitting code & software for further information.
Data	
- Accession codes, un - A list of figures that	ut <u>availability of data</u> include a <u>data availability statement</u> . This statement should provide the following information, where applicable: ique identifiers, or web links for publicly available datasets have associated raw data restrictions on data availability
All relevant data collecte	d, analyzed and shown in the figures, as well as input dataset and code for SCA analysis are available in source data.
Field-speci	fic reporting
Please select the one b	relow that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.
x Life sciences	Behavioural & social sciences Ecological, evolutionary & environmental sciences
For a reference copy of the de	ocument with all sections, see <u>nature.com/documents/nr-reporting-summary-flat.pdf</u>

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Lite	sciences	stud	V C	lesign

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All studies must dis	sclose on these points even when the disclosure is negative.			
Sample size	Each electrophysiology experiment was performed on at least 3 individual cells. The number of recordings of each experiment was based on personal experience and data variability.			
Data exclusions	Electrophysiology recordings displaying high noises, instability, or currents were contaminated by endogenous currents from oocytes, for example at very positive or negative voltages, were excluded.			
Replication	All experiments were successfully reproduced.			
Randomization	The experiments were not randomized.			
Blinding	The investigators were not blinded to allocation during experiments and outcome assessment.			
Reportin	g for specific materials, systems and methods			
,	on from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, ted is relevant to your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.			
Materials & ex	perimental systems Methods			
n/a Involved in th	n/a Involved in the study			
Antibodies	ChIP-seq			
x Eukaryotic	cell lines Flow cytometry			
Palaeontol				
=1=	d other organisms			
	earch participants			
∡ Clinical dat	.d			
Antibodies				
Antibodies used	For KCNQ1 (primary antibody conjugated with HRP, do not need secondary antibody) Name: KCNQ1 (G-8) mouse monoclonal IgG Cat. No.: sc-365186 Vendor: Santa Cruz.			
	For Gβ, Primary antibody, Name: Gβ (T-20) rabbit polyclonal IgG Cat. No.: sc-378 Vendor: Santa Cruz. Secondary antibody Name: Goat anti-Rabbit IgG (H+L) Secondary Antibody, HRP Cat. No.: A16110 Vendor: Thermo			
Validation	Both KCNQ1 and G β bands are consistent with those provided from the vendor.			
Animals and	other organisms			
Policy information about studies involving animals; ARRIVE guidelines recommended for reporting animal research				
Laboratory anima	als Xenopus laevis			

Wild animals n/a Stage V or VI oocytes were obtained from Xenopus laevis by laparotomy. All procedures were approved by the Washington Field-collected samples

University Animal Studies Committee. In details, frogs werer housed in a professional animal facility in the basement of Whitaker Hall downstairs from the laboratory where experiments are performed. Frogs were cared for by the Institutional Animal Care and Use Committee Office at Washington University. Oocytes removal operations were performed in animal preparation rooms in the animal facility. Technical assistance will be provided by the animal facility.

Ethics oversight All procedures were performed in accordance with the protocol approved by the Washington University Animal Studies Committee (Protocol # 20190030)

Note that full information on the approval of the study protocol must also be provided in the manuscript.